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piece 1, NC\_000913, TERM31\_yecR-, config: linear, direction: -, begin: 1986275, end: 1985878

**p35** 1.8 bi

... p35-(22)-p10 1986100 Gap 2.3 bits [###> orf 2 codons [###> orf 6 codons ...

... -----| p35-p10 1986100 total 5.1 bits  ir TERM31\_yecR-

$\mu$  p35 5.5 bits

{-----} sd-(6)-ir 1986082 Gap 4.3 bits {-----} sd-(14)-ir 1986047 Gap 4.9 bits

{-----} sd-(8)-ir 1986080 Gap 2.4 bits |-----| sd-ir 1986047 TERM31\_vecR total

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|-----| sd-ir 1986080 TERM31_yecR- total 6.2 bits |-----| p35-(22)-|-----| p35-d10 1 |-----| {-----| p35-----| p35-----| ..
```

The diagram illustrates the p10 promoter region with the following features:

- [###> orf 6 codons**: Located at the top left.
- p10 2.9 bits**: The first transcription start site, indicated by a red arrow pointing right.
- sd**: The sigma factor binding site, shown as a green box below the DNA sequence.
- ir TERM31\_yecR-**: An inverted repeat sequence located downstream of the p10 promoter.
- p10**: The second transcription start site, indicated by a red arrow pointing right.
- ... p10 7.1 bits**: A third transcription start site, indicated by a red arrow pointing right.
- {-----} sd-(10)-ir 1985983 Gap 2.7 bits**: A regulatory element consisting of a sigma factor binding site (green), a 10 bp gap, and an inverted repeat sequence (red).

... p10 3.3 bits |-----| sd-ir 1985983 TERM31\_yecR- total 6.3 bits

 p35 3.3 bits

